

<400> 2

Sequence PC19084 12192003.ST25.txt SEQUENCE LISTING

<110> Agouron Pharmaceuticals, Inc./A Pfizer Company

<120> PIN1 PEPTIDYL-PROLYL ISOMERASE POLYPEPTIDES, THEIR CRYSTAL STRUCTURES, AND
USE THEREOF FOR DRUG DESIGN

```
PC19084A
<130>
       10/616,003
<140>
       2003-07-08
<141>
       us 60/394,889
<150>
<151>
       2002-07-09
<160> 11
<170> PatentIn version 3.1
<210>
       423
<211>
<212>
       DNA
       Artificial
<213>
<220>
       pplase
<223>
<400>
atgggcagca gccatcatca tcatcatcac agcagcggcc tggtgccgcg cggcagccat
                                                                         60
atgggcaaaa acgggcaggg ggagcctgcc agggtccgct gctcgcacct gctggtgaag
                                                                       120
cacagccagt cacggcggcc ctcgtcctgg cggcaggaga agatcacccg gaccaaggag
                                                                        180
gaggccctgg agctgatcaa cggctacatc cagaagatca agtcgggaga ggaggacttt
                                                                        240
                                                                        300
gagtctctgg cctcacagtt cagcgactgc agctcagcca aggccagggg agacctgggt
                                                                        360
gccttcagca gaggtcagat gcagaagcca tttgaagacg cctcgtttgc gctgcggacg
                                                                        420
ggggagatga gcgggcccgt gttcacggat tccggcatcc acatcatcct ccgcactgag
                                                                        423
tga
<210>
       2
       123
<211>
<212>
       PRT
       Artificial
 <213>
<220>
<223>
       PPlase
```

Gly Ser His Met Gly Lys Asn Gly Gln Gly Glu Pro Ala Arg Val Arg
1 10 15

Cys Ser His Leu Leu Val Lys His Ser Gln Ser Arg Arg Pro Ser Ser 20 25 30

Sequence PC19084 12192003.ST25.txt Trp Arg Gln Glu Lys Ile Thr Arg Thr Lys Glu Glu Ala Leu Glu Leu Ile Asn Gly Tyr Ile Gln Lys Ile Lys Ser Gly Glu Glu Asp Phe Glu 50 60 Ser Leu Ala Ser Gln Phe Ser Asp Cys Ser Ser Ala Lys Ala Arg Gly 65 70 75 80 Asp Leu Gly Ala Phe Ser Arg Gly Gln Met Gln Lys Pro Phe Glu Asp Ala Ser Phe Ala Leu Arg Thr Gly Glu Met Ser Gly Pro Val Phe Thr Asp Ser Gly Ile His Ile Ile Leu Arg Thr Glu 115 120 <210> 422 <211> <212> DNA Artificial <220> <223> **PPlase** <400> atgggcagca gccatcatca tcatcatcac agcagcggcc tggtgccgcg cggcagccat 60 atggcaaaaa cgggcagggg gagcctgcca gggtccgctg ctcgcacctg ctggtgaagc 120 180 acagccagtc acggcggccc tcgtcctggc ggcaggagca gatcacccgg acccaggagg aggccctgga gctgatcaac ggctacatcc agaagatcaa gtcgggagag gaggactttg 240 300 agtctctggc ctcacagttc agcgactgca gctcagccaa ggccagggga gacctgggtg ccttcagcag aggtcagatg cagaagccat ttgaagacgc ctcgtttgcg ctgcggacgg 360 gggagatgag cgggcccgtg ttcacggatt ccggcatcca catcatcctc cgcactgagt 420 422 ga <210> 123 <212> PRT <213> Artificial <220> <223> **PPlase** <400> Gly Ser His Met Gly Lys Asn Gly Gln Gly Glu Pro Ala Arg Val Arg 1 10 15

Sequence PC19084 12192003.ST25.txt Cys Ser His Leu Leu Val Lys His Ser Gln Ser Arg Arg Pro Ser Ser 20 25 30 Trp Arg Gln Glu Gln Ile Thr Arg Thr Gln Glu Glu Ala Leu Glu Leu 35 40 45 Ile Asn Gly Tyr Ile Gln Lys Ile Lys Ser Gly Glu Glu Asp Phe Glu 50 60Ser Leu Ala Ser Gln Phe Ser Asp Cys Ser Ser Ala Lys Ala Arg Gly 65 70 75 80 Asp Leu Gly Ala Phe Ser Arg Gly Gln Met Gln Lys Pro Phe Glu Asp Ala Ser Phe Ala Leu Arg Thr Gly Glu Met Ser Gly Pro Val Phe Thr Asp Ser Gly Ile His Ile Ile Leu Arg Thr Glu 115 120 <210> <211> 36 <212> DNA Artificial <213> <220> <223> Primer <400> 36 agcagccata tgggcaaaaa cgggcagggg gagcct <210> <211> 30 <212> DNA Artificial <213> <220> Primer <223> <400> 6 30 cttggatcct cactcagtgc ggaggatgat <210> 7 119 <211> <212> PRT <213> Artificial

Page 3

<220> <223>

<400>

7

PPlase domain

```
Sequence PC19084 12192003.ST25.txt
Gly Lys Asn Gly Gln Gly Glu Pro Ala Arg Val Arg Cys Ser His Leu
1 10 15
Leu Val Lys His Ser Gln Ser Arg Arg Pro Ser Ser Trp Arg Gln Glu 20 25 30
Lys Ile Thr Arg Thr Lys Glu Glu Ala Leu Glu Leu Ile Asn Gly Tyr
35 40
Ile Gln Lys Ile Lys Ser Gly Glu Glu Asp Phe Glu Ser Leu Ala Ser 50 60
Gln Phe Ser Asp Cys Ser Ser Ala Lys Ala Arg Gly Asp Leu Gly Ala 65 70 75 80
Phe Ser Arg Gly Gln Met Gln Lys Pro Phe Glu Asp Ala Ser Phe Ala
Leu Arg Thr Gly Glu Met Ser Gly Pro Val Phe Thr Asp Ser Gly Ile 100 \hspace{1cm} 105 \hspace{1cm} 110
His Ile Ile Leu Arg Thr Glu
<210>
       8
<211>
        44
<212>
       DNA
       Artificial
<213>
<220>
<223>
       Primer
<400> 8
                                                                                44
gcggcaggag cagatcaccc ggacccagga ggaggccctg gagc
        9
<210>
<211>
       44
<212>
       DNA
       Artificial
<213>
<220>
<223>
        Primer
                                                                                44
gctccagggc ctcctcctgg gtccgggtga tctgctcctg ccgc
<210>
        10
        119
<211>
<212>
       PRT
       Artificial
<213>
<220>
<223> PPlase domain
```

Page 4

Sequence PC19084 12192003.ST25.txt

<400> 10

Gly Lys Asn Gly Gln Gly Glu Pro Ala Arg Val Arg Cys Ser His Leu $10 \,$ 15

Leu Val Lys His Ser Gln Ser Arg Arg Pro Ser Ser Trp Arg Gln Glu 20 25 30

Gln Ile Thr Arg Thr Gln Glu Glu Ala Leu Glu Leu Ile Asn Gly Tyr 35 40 45

Ile Gln Lys Ile Lys Ser Gly Glu Glu Asp Phe Glu Ser Leu Ala Ser 50 55 60

Gln Phe Ser Asp Cys Ser Ser Ala Lys Ala Arg Gly Asp Leu Gly Ala 65 70 75 80

Phe Ser Arg Gly Gln Met Gln Lys Pro Phe Glu Asp Ala Ser Phe Ala 85 90 95

Leu Arg Thr Gly Glu Met Ser Gly Pro Val Phe Thr Asp Ser Gly Ile $100 \hspace{1cm} 105 \hspace{1cm} 110$

His Ile Ile Leu Arg Thr Glu 115

<210> 11

<211> 11 <212> PRT

<213> Artificial

<220>

<223> Pintide where the serine is a phosphorylated

<400> 11

Phe Leu Trp Phe Tyr Pro Ser Pro Phe Leu Glu 1 5 10